AUSTRALIAN PATENT OF. ICE



WRITTEN OPINION

	[i	Date of mailing		
		day/month/year	0,2 JUN 2303	
Applicant's or agent's file reference		REPLY DUE within FIVE MONTHS of the date of the Registrar's letter enclosing the written opinion		
200130790/030423/TMSR/3220			Priority Date (day/month/year)	
Application No.	Application Filing Dat	e (day/month/year)		
SG 200103079-1	22 May 2001		17 July 2000	
International Patent Classification (IPC) (as	s indicated in the search	report)		
Int. Cl. 7 C12Q 1/68	?			
Applicant		යි		
WANG, XIAO BING et al) 	
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		-9		
1. This first written opinion consists of	a total of 5 sheets.	ن ن	}	
2. This opinion contains indications relat	ting to the following item			
I X Basis of the opinion	Ş		and a state of the	
II Non-establishment of o	pinion with regard to no	velty, inventive step	and industrial applicability	
III Lack of unity of invent	ion			
IV X Reasoned statement wincitations and explanation	ith regard to novelty, invo	entive step or industr ement	aal applicability;	
V Certain documents cite	ed .			
VI Certain defects in the a	application			
VII X Certain observations o	n the application			
3. This opinion is based upon the assum				
4. The search report used was issued by	the Australian Office	, and the date of com	pletion is: 28 May 2003	
5. If no reply is filed, the examination re	eport will be established	on the basis of this o	pinion.	
6. The date by which the examination re	eport will be established	is: 17 October 2004		
Name and mailing address		Authorized Officer		
AUSTRALIAN PATENT OFFICE				
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E-mail address: pct@ipaustralia.gov.au Facsimile no. 61 2 62853929	E-mail address: pct@ipaustralia.gov.au Facsimile no. 61 2 62853929 TERRY MOORE			

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I. Basis of the opinion	on		
1. This opinion has been drawn on the basis of:			
X the application as	s originally filed.		
the description,	pages , as originally filed,		
	pages , filed with the request,		
	pages , received on with the letter of		
the claims,	pages , as originally filed,		
	pages , filed with the request,		
	pages , received on with the letter of		
the drawings,	sheets/fig. , as originally filed,		
	sheets/fig. , filed with the request,		
	sheets/fig. , received on with the letters of		
the sequence list	ting part of the description:		
	pages , as originally filed		
	pages , filed with the demand pages , received on with the letter of		
a The second boson	resulted in the cancellation of: pages:		
2. The amendments have	sheets of drawings/figures No:		
3 This opinion ha	s been established as if (some of) the amendments had not been made, since they have been considered to lisclosure as filed, as indicated in the Supplemental Box.		
4. Additional observation	ns, if necessary:		
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	īv.	Reasoned statement with regard to novelty, inventive step or industrial applicability; citations and explanations
١		supporting such statement

supporting such statement			
1. Statement			
Novelty (N)	Claims 19-22 and 28-30	YES	
	Claims 1-18, 23-27 and 31-36	NO `	
Inventive step (IS)	Claims	YES	
	Claims 1-36	NO	
Industrial applicability (IA)	Claims 1-36	YES	
industrial approximity (= 1)	Claims	NO	

2. Citations and explanations

The invention described in the specification resides in a method of detecting or quantifying a target nucleic acid sequence with respect to a specific base change in the sequence. The method broadly involves the use of primer that is complementary to the target sequence and that anneals to the target sequence immediately upstream of the specific base. Primer extension is then conducted using a mix comprising:

- one type of ddNTP, or an absence of the any nucleotide complementary to the specific base, and (i)
- the remaining three types of dNTPs that are different to the complement of the specific base, any of (ii) which may be optionally labelled.

Novelty and Inventive Step

The following documents identified in the International Search Report have been considered for the purposes of this report:

- WO 96 30545 D1
- Braun et al D2
- US 5 888 819 D3
- WO 91 13075 D4
- Prezant et al D5
- Piggee et al D6

D1 discloses a method for detecting or quantifying a target nucleic acid sequence with respect to base changes at a specific location. In particular the method is used to assess mutations in the human COX1 gene, but it is also discloses that the method is suitable for a broad range of nucleic acids. D1 discloses use of a primer that whose 3' end is immediately adjacent to the base of interest. The primer is extended using a polymerase and a mix comprising one type of ddNTP, or an absence of nucleotide corresponding to the complement of the base of interest and from one to three of the remaining three types of nucleotide that are different to the complement of the specific base. The ddNTP and/or the dNTPs may optionally be labelled. As such the citation deprives claims 1-18, 23, 24, 27 and 31-36 of novelty.

With respect to the remaining claims, features such as those that enable attachment to a solid support, use of nucleotide analogues and extragenomic samples simply represent routine applications of the disclosed method that are standard in the art. As such these claims lack an inventive step.

Continued on supplemental sheet

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VII. Certain observations on the application				
The following observations on the clarity of the claims, descripti supported by the description, are made:	on, and drawings or on the question whe	ther the claims are fully		
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*				
X The claimed invention is patentable according to Section 1	3(2); or			
The claimed invention is unpatentable according to Section	n 13(2) because:			
N.				

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Sı	uр	pler	nent	al	Box
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(To be used when the space in any of Boxes I to VII is not sufficient)

Continuation of Box [No.]: IV2

D2 discloses a further method using a primer that anneals immediately adjacent to the base of interest and primer extension using a mix comprising the ddNTP corresponding to the complement of the base interest and three dNTPs that are different to the complement of the base of interest. The method is used to assess mutations in the CFTR gene.

In particular figure 1(b) discloses the F508C mutation and the use of ddCTP as the chain terminator and 1(c) discloses the G542X mutation using ddTTP. The citation also discloses affinity capture of PCR products on solid phase and the use of modified nucleoside analogues. As such the citation deprives claims 1-6, 10-13, 23-27 and 31-36 of novelty.

Furthermore, as discussed with respect to D1, the subject matter of the remaining claims appears to represent nothing more than routine application of the method disclosed in the citation. As such, the remaining claims lack an inventive step in light of D2

D3-D6 all disclose similar methods to those disclosed in the specification. However D3 and D6 use only ddNTPs corresponding to the base of interest, with no added dNTPs and D4 and D5 disclose only dNTPS, with no added ddNTPs. As such none of D3-D6 appear to disclose or teach toward the subject matter of the claims.